SEQUENCE LISTING

<110>   Baehrecke, Eric H.

<120>   GENES REGULATING PROGRAMMED CELL DEATH

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<140>   10/016,768
<141>   2001-10-29

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<170>   PatentIn version 3.2

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Page 10

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Ser His Leu Asn Ile Glu Glu Gln Gln Pro Ile Ala Ile Ala Gly Ser
            35                  40                  45


Glu Asp Glu Pro Ser Gln Tyr Asn His Ser Ser Lys Glu Ile Ser Gln
        50                  55                  60


Ser Asn Pro Asn His Cys Lys Thr Glu Asn His Arg Leu Glu Gln Gln
65                  70                  75                  80


His Asn Gly Ser Gln Leu Leu Glu Glu Glu Asp Ser Glu Asn Asn Gln
                85                  90                  95


Thr Ser His Asp Ser Ser Arg Thr Pro Thr Pro Gly Ala Thr Ser Thr
                100                 105                 110


Pro Ser Pro Pro Pro Glu Pro Ile Asp Trp Arg Pro Ser Ala Lys Cys
            115                 120                 125


Asn Phe Cys Val Asn Gly Arg Leu Leu Thr Val Asn Ala Gln Gly Lys
    130                 135                 140


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Ser His Ile His Gln His Asp Ser Asp Ser Asn Ser Ser Ala Ser Leu
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Page 16

Pro His His Ile Ser Ser Ser Ser Ser Ser Asn Asn Asn Ser Ser Gly
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Ala Ala Thr Pro Ala Asn Ser Leu Glu Leu Tyr Lys Leu Leu Thr Gln
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Arg Ala Ala Lys Met Thr Ser Met Asp Ser Met Ala Ala Gln Leu Ala
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Thr Pro Ser Pro Ser Val Asp Ala Pro Leu Asp Leu Ser Ser Lys Pro
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Ser Ala Ser Gln His His Glu Gln Arg Ser Ile Leu Asp Asn Arg Leu
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Phe Lys Met Lys His His Asp Gln Glu Gln Asp His Asp Gly Asp Glu
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Leu Glu Asp Ser Asn Asp Asp Ala Glu Ala Glu Val Asp Ser Asn Ala
385   390   395   400

Ser Thr Pro Val Tyr Pro Ala Glu Phe Ala Arg Ala Gln Leu Arg Lys
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Leu Ser His Leu Ser Glu His Asn Gly Ser Asp Leu Gly Glu Asp Val
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Asp Arg Gly Ser Pro Lys Met Gly Arg His Pro Ala Cys Gly Asn Ala
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Ser Ala Asn Gln Gly Ala Pro Ala Ser Ile Pro Leu Asp Ala Asn Val
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Leu Leu His Thr Leu Met Leu Ala Ala Gly Ile Gly Ala Met Pro Lys
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Leu Asp Glu Thr Gln Thr Val Gly Asp Phe Ile Lys Gly Leu Leu Val
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Ile Leu Lys Ile Pro Ser Phe Lys Val Ser Gly Pro Ala Ser Ser Ser
        580   585   590

Ser Leu Ser Pro Gly Gly Leu Val Gly Gly His His His Pro Leu Asn
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Ser His Arg Asn Gly Ser Asn Arg Ser Pro His Ser Ala Ser Pro Met
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Leu Thr Ser Ser Ser Ser Ser Ile Gln Lys Met Met Ala Ser Asn Ile
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Thr Ser Arg Phe Gly Ala Ser Pro Asn Leu Leu Ser Gln Gln His His
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                805             810             815

Lys Arg Glu Pro Lys Pro Gln Pro Asp Leu Val Gly Leu Thr Gly Pro
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Ala Asn Lys Leu Gln Leu Asp Lys Leu Lys Ala Gly Pro His Gly Gly
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Ser Lys Leu Ser Asn Ala Leu Lys Asn Gln Asn Asn Gln Ala Ala Ala
    850             855             860

Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Ala Thr Pro Asn Gly
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Leu Lys Leu Pro Leu Phe Glu Ala Gly Pro Gln Ala Leu Ser Phe Gln
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Gly His Gly His Gly His Ala Leu Leu Asp Gln Leu Leu Val Lys Lys
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Cys Ser Ser Ala Ser Gly Glu Ser Val Lys Arg Ser Gly Ser Pro
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Ser Ser Arg Met Thr Ser Arg Asp Asp Ser Glu Thr Asp Ala Ser
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Ser Phe Lys Ser Gly Glu Asn Gly Gly Gln Gln Asn His Lys Met
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Ser His Leu Asn Ile Glu Glu Gln Gln Pro Ile Ala Ile Ala Gly Ser
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Glu Asp Glu Pro Ser Gln Tyr Asn His Ser Ser Lys Glu Ile Ser Gln
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Ser Asn Pro Asn His Cys Lys Thr Glu Asn His Arg Leu Glu Gln Gln
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His Asn Gly Ser Gln Leu Leu Glu Glu Glu Asp Ser Glu Asn Asn Gln
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Thr Ser His Asp Ser Ser Arg Thr Pro Thr Pro Gly Ala Thr Ser Thr
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Pro Ser Pro Pro Pro Glu Pro Ile Asp Trp Arg Pro Ser Ala Lys Cys
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Asn Phe Cys Val Asn Gly Arg Leu Leu Thr Val Asn Ala Gln Gly Lys
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Leu Val Ala Glu Ser Ala Ala Thr Ala Thr Ser Ser Ser Thr Ser Asn
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Ser His Ile His Gln His Asp Ser Asp Ser Asn Ser Ser Ala Ser Leu
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Pro His His Ile Ser Ser Ser Ser Ser Asn Asn Asn Ser Ser Gly
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Asn Arg Ala Arg His Ile Ala Ala Ala Ser Ala Arg Ala Thr Pro Ala
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Ala Ala Thr Pro Ala Asn Ser Leu Glu Leu Tyr Lys Leu Leu Thr Gln
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Arg Ala Ala Lys Met Thr Ser Met Asp Ser Met Ala Ala Gln Leu Ala
225   230   235   240

Gln Phe Ser Leu Leu Ala Asp Phe Asn Leu Ile Asn Ser Leu Ala Ser
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Gln Gln Gln Gln Gln Gln Gln Gln Gln Ile Ala Ser Ala Val Thr Pro
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Thr Thr Ser Glu Val Ser Ala Ala Ala Ile Ser Pro Ala Leu Lys Asp
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Thr Pro Ser Pro Ser Val Asp Ala Pro Leu Asp Leu Ser Ser Lys Pro
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Ser Pro Asn Ser Ser Ile Ser Gly Asp Val Lys Ser Val Arg Ala Cys
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Ala Thr Pro Thr Pro Ser Gly Arg Arg Ala Tyr Ser Glu Glu Asp Leu
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Ser Arg Ala Leu Gln Asp Val Val Ala Asn Lys Leu Asp Ala Arg Lys
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Ser Ala Ser Gln His His Glu Gln Arg Ser Ile Leu Asp Asn Arg Leu
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Phe Lys Met Lys His His Asp Gln Glu Gln Asp His Asp Gly Asp Glu
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Leu Glu Asp Ser Asn Asp Asp Ala Glu Ala Glu Val Asp Ser Asn Ala
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Ser Thr Pro Val Tyr Pro Ala Glu Phe Ala Arg Ala Gln Leu Arg Lys
                405   410   415

Leu Ser His Leu Ser Glu His Asn Gly Ser Asp Leu Gly Glu Asp Val
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Asp Arg Gly Ser Pro Lys Met Gly Arg His Pro Ala Cys Gly Asn Ala
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Ser Ala Asn Gln Gly Ala Pro Ala Ser Ile Pro Leu Asp Ala Asn Val
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Leu Leu His Thr Leu Met Leu Ala Ala Gly Ile Gly Ala Met Pro Lys
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Ile Leu Lys Ile Pro Ser Phe Lys Val Ser Gly Pro Ala Ser Ser Ser
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Tyr Lys Lys Pro Ser Ile Ser Val Ala Lys Ile Ile Gly Gly Thr Asp
705   710   715   720

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Glu His Val Leu Ala Pro Asp Glu Arg His Glu Cys Leu Arg Pro Gly
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Leu Gln Ser His His Gly Gly Asp Ala Glu Phe Pro Gly Leu Gln Ser
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Pro Arg Leu Met Lys Ser Ala Gln Asp Met Val Gly Glu Arg Leu Arg
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Trp His His Gln Glu Asp Ala Ala Gly Glu Gln Gly Asn Gly Ser Ala
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Ala Gly Asn Gly Ser Asn Gly Ser Asn Gly Asn Gly His Gly His Gly
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Lys Ser Gly Glu Asn Gly Gly Gln Gln Asn His Lys Met Met Asp
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tactcttagc cacataaccg atacgataca atagccactt aattaggatc gatcatagcc      9000

aagcaataga atcagatatc agatacaaaa ctcagaaccg gaaacaggaa atcgtcaatc      9060
```

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gcgaatcgga aaaaagaagc agcaccaaat cgaactgcaa aggcaaaacc ccagtatatt    9120

aataatgggg agaaacataa tgataaacca tttttatttc aattattaac ttataacaca    9180

acatcaccac cacagcttcc cacagtttca gttacacagg accaccacca tagccctagg    9240

aaattatatt cataattaat caatcaatca atccgtaaaa ccaacaactt caatagttat    9300

aagcaattgg cgtcaaaaaa aaaaaagcg aaaacaacca aaatcttagc caggcaaaag    9360

ttttgagcat attttttcatt atttttacaa caaacaaaca aactgatgta agtacaattc    9420

ataaaattag actttcggta aactataata aagaaaatag agcagaaaat aacatttttt    9480

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aaaagccaat aagcaaataa aaagaat                                        9567
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